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MPsrch\_pp protein - protein database search, using Smith-Waterman algorithm

Tabular output not generated. Run on: Sat May 13 09:43:25 2000; MasPar time 4.90 Seconds 333.332 Million cell updates/sec

Title:
Description:
Perfect Score:
Sequence:

>US-09-331-631-1 (117-185) from US09331631.pep (4 of 5) 539 1 NRQRDPQQQYEQCQKHCQRR......ЕEQQREDEEKYEERMKEEDN 69

Scoring table: PAM 150 Gap 11

Searched: 188963 seqs, 23686106 residues

Post-processing: Minimum Match 0% Listing first 45 summaries

Database: a-geneseq35 1:geneseqp

Statistics: Mean 25.022; Variance 122.235; scale 0.205

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

Result No.	Score	Query Match	Length	DB	ID	Description	Pred. No.
	539	100.0	666	<u>-                                    </u>	W62828	Macadamia integrifolia	2.16e-36
.,		96.5	625	ᆫ	W62830	'n	9.12e-35
•			666	<u>_</u>	W62829		1.65e-34
4			525	_	W62831	Theobroma cacao antimi	2.27e-09
			566	_	R20181	Ø	2.27e-09
6		37.3	590	<u>,,,</u>	W62832	m hirsutum a	4.42e-08
~1			444	ب	W90340	G. max truncated SBP1	1.90e-02
<b>~</b>			524	ר	W90339	G. max SBP1 protein.	1.90e-02
.0		22.6	593	1	W62835	Zea mays antimicrobial	6.38e-02
10		21.0	33	<b></b>	W62836	Zea mays antimicrobial	2.98e-01
13		21.0	409	μ.	W90342	G. max truncated SBP2	2.98e-01
1			489	Н	W90341	G. max SBP2 protein.	2.98e-01
11		19.7	316	<b>ب</b>	R26941	P.falciparum LSA-R-NR	9.71e-01
1.		19.7	493	Ь	R26944	P.falciparum LSA gene	9.71e-01
11		19.5	28	μ	W62841		1.15e+00
16	104	19.3	35	_	R21079	Antimicrobial maize pe	1.36e+00
-		19.1	303	L	R60054	Dirofilaria immitis pa	1.60e+00
. 18		18.7	395	L	W03474	Mouse SRY-related prot	2.24e+00
19		17.8	433	ب	W05389	Mouse SH3P7 protein.	5.11e+00
20		17.4	265	<b>-</b>	R12844	HTLV-1 protein express	7.09e+00
21		17.4	450	H	W46606	Tyrosine kinase associ	7.09e+00
2		17.4	559	$\vdash$	R05427	Circumsporozoite (CS)-	7.09e+00
2	93	17.3	432	ш	W93954	Human regulatory molec	8.34e+00

44	43	42	41	40	39	3 8	37	36	35	34	ω ω	32	31	30	29	28	27	26	25	24	
8 g	87	87	88	88	89	89	89	89	89	89	90	91	91	91	91	91	91	92	93	93	
16.0	16.1																				
3/6 796	462	388	614	436	1752	1312	1312	1088	301	83	482	802	787	783	765	765	541	562	740	740	
<u></u>	ــر د	Ъ	س	ب	Н	Ц	Н	Н	ш	ш	ш	ᆫ	<u></u>	<u>_</u>	<b>⊢</b>	ᆫ	H	Н	<b>ب</b>	ب	
W36024	R05766	Y04998	R82630	W03662	Y07031	W71295	W22775	W19786	P70867	R90546	Y07067	W37153	W37152	W37151	P92275	R75915	W37148	R70491	R68838	R27530	
Sequence of protein H. MSRV-1 pol protein seq	eptide a	Mycobacterium species	70K autoantigen, part	Human 70K U1 snRNP pro	SSOC	Human homologue of yea	Human RAD50.	Human multiple regulat	Sequence of acidic bas	pJG4-5-CDK-BP clone de			Mouse neural Mena++ pr	Mouse neural Mena+ pro	Human topoisomerase I	Variant human DNA topo	Mammalian Ena (Mena).	Leucocytozoan protozoa	_	Plasmodium falciparum	
2.58e+01		2.20e+01	1.87e+0	1.87e+0	1.60e+01	1.60e+0	1.60e+0	1.60e+0:	1.60e+0:	1.60e+0	1.36e+01	1.15e+0	1.15e+01	1.15e+01	1.15e+01	1.15e+01	1.15e+01	9.82e+00		8.34e+00	

## ALIGNMENTS

RESULT ID W AC W DT 2 DE M KW a	Oy Oy	X B O		RESULT ID W AC W DT 2 DE M KW a
JLT 2 W62830 standard; Protein; 625 AA. W62830; W62830; 27-OCT-1998 (first entry) Macadamia integrifolia antimicrobial protein. antimicrobial protein; infestation; control.	117 NRQRDPQQQYEQCQKHCQRRETEPRHMQTCQQRCERRYEKEKRKQQKRYEEQQREDEEKY 176	Query Match 100.0%; Score 539; DB 1; Length 666; Best Local Similarity 100.0%; Pred. No. 2.16e-36; Matches 69; Conservative 0; Mismatches 0; Indels 0; Gaps	Rey Location/Qualifiers Peptide 1. 28 /note= "signal peptide" Protein 29. 666 Protein /note= "mature protein"  W09827805-Al. 02-JUL-1998. 22-DEC-1996; AU-004275. 22-DEC-1996; AU-004275. (RETR-) COOP RES CENT TROPICAL PLANT PATHOLOGY. Bower NI, Goulter KC, Green JL, Manners JM, Marcus JP; WPI; 98-377279/32. N-PSDB; V42310. Novel anti-microbial protein from e.g. Macadamia integrifolia - useful for controlling microbial infestations of plants or mammals Claim 1; Page 34-36; 96pp; English. The sequence is that of an antimicrobial protein which can be used to control microbial infestations in plants and mammalian animals. Sequence 666 AA:	1 62828 standard; 62828; 7-0CT-1998 (fi acadamia integr ntimicrobial pr acadamia integr

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22-DEC-1997; AU0874.
20-DEC-1996; AU-004275.
(RETR-) COOP RES CENT TR
Bower NI, Goulter KC, Gr
WPI; 98-377279/32.
                                                                                                                                                                                                                                                                                                                                                                                                                               Novel anti-microbial protein from e.g. Macadamia integrifolia useful for controlling microbial infestations of plants or man Claim 1, Page 39-41; 96pp; English.

The sequence is that of an antimicrobial protein which can be used to control microbial infestations in plants and mammal
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The sequence is that of an antimicrobial protein which can be used to control microbial infestations in plants and mammalian
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Macadamia integrifolia antimicrobial protein.
antimicrobial protein; infestation; control.
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                                  EERMKEEDN 185
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                                                                                                                                                                                                                                                                         Similarity
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29. .666
/note= "mature proto:-"
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95.7%;
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95.78;
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Green JL, Manners JM, Marcus JP;
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Pred. No. 9.12e-35;
1; Mismatches 2
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No. 1.
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pp beans and produced in large quantities using yeast and bacterial pression vectors

CC The inventors claim a 67 kD and 31 kD T. cacao protein, and cc fragments, and encoding DNAs. The 47 kD and 31 kD proteins are CC derived from the 67 kD precursor. T. cacao protein cDNA was CC detected in a cDNA library prepared from immature cocoa beans RNA CC using a probe based on the AA sequence of a CNBr peptide common to the 47 kD and 31 kD polypeptides. Homology searches revealed close CC the 47 kD and 31 kD polypeptides. Homology searches revealed close CC complete the 67 kD polypeptide and the vicilins, which are CC seed storage proteins.
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PARENTANT

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Best Local
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(MRSC ) MARS UK LID.

Spencer ME, Hodge R, Deakin EA, Ashton

WPI; 92-024418/03.
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R20181;
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The sequence is that of an antimicrobial protein which can be used to control microbial infestations in plants and man be used to control microbial infestations in plants.
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11-JUN-1990; GB-0130
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llarity 47.0%;
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  ne 217; DB 1; 1
1. No. 2.27e-09;
Mismatches 16
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WPI: 99-070155/06.

New modified plant sucrose binding proteins - used to develop transgenic plants which can have enhanced or decreased sucrose uptake activity in developing seeds

Claim 7: Page 36-37: 58pp; English.

This sequence represents a novel sucrose binding protein, SBPl isolated from Glycine max. This protein is used in a method resulting in the production of a modified plant sucrose binding protein (SBP) which has a modified amino acid sequence compared to a corresponding wild-type SBP, and where expression of the modified SBP in a yeast assay system confers enhanced sucrose uptake compared to the corresponding wild-type SBP.

The products of the invention can be used for producing transgenic plants which have modified sucrose uptake activity, particularly in developing transgenic plants.
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The sequence is that of an antimicrobial protein which can mark the sequence is that of an antimicrobial protein which can mark the sequence is that of an antimicrobial protein which can mark the sequence is that of an antimicrobial infestations in plants and mark the sequence of t
     seeds. Enhanced sucrose uptake activity in developing seeds may be desirable where it is an advantage to increase the carbohydrate content of the seed (e.g. where the seed is the primary plant material harvested such as soybean). In contrast, decreased sucrose uptake activity in
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22-DEC-1997; AU0874.
20-DEC-1996; AU-004275.
(RETR-) COOP RES CENT TROPICAL PLANT PATHOLOGY.
BOWER NI, GOLILEY KC, Green JL, Manners JM, Mar
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G. max truncated SBP1 protein.
SBP1; sucrose binding protein;
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(UNIW) UNIV WASHINGTON STATE RES FOUND
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            37.3%;
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Pred. No. 4.42e-08;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SBP2; sucrose uptake; transgenic plant;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Manners JM, Marcus JP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 590
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plants or man
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02-JUL-1998. 22-DEC-1997; 20-DEC-1996;

COOP

TROPICAL

PLANT PATHOLOGY

AU0874. AU-004275. RES CENT T

Zea mays. WO9827805-A1.

antimicrobial protein;

protein.
infestation;

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Best Local S
Matches 1
                                                                                                                                                                                                                                                                                               Best
                                                                                                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                                                                                                                 Query Match
                                                                                             W62835;
27-OCT-1998 (first entry)
7-- mays antimicrobial pro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     This sequence represents a novel sucrose binding protein, SBP1 isolated from Glycine max. This protein is used in a method resulting in the production of a modified plant sucrose binding protein (SBP) which has a modified amino acid sequence compared to a corresponding wild-type SBP, and where expression of the modified SBP in a yeast assay system confers enhanced sucrose uptake compared to the corresponding wild-type SBP. The products of the invention can be used for producing transgenic plants
                                                                                                                                                                                                                                                                                                                                                                                           of the seed (e.g. where the seed is the primary plant material harvested such as soybean). In contrast, decreased sucrose uptake activity in seeds might be desirable where the vegetative material of the plant is harvested. The SBP regulatory regions confer specific or enhanced
                                                                                                                                                        W62835
                                                                                                                                                                                                                                                                                                                                                                                                                                                         which have modified sucrose uptake activity, particularly in developing seeds. Enhanced sucrose uptake activity in developing seeds may be desirable where it is an advantage to increase the carbohydrate content
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New modified plant sucrose binding proteins - used to develop transgenic plants which can have enhanced or decreased sucros
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Chao ws,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          seed; carbohydrate content; soybean.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SBP1; sucrose binding protein; SBP2; sucrose uptake; transgenic plant;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         G. max SBP1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /T 8
W90339 standard; protein; 524
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              transgene in developing seeds. Sequence, 444 AA;
                                                                                                                                                                                                                                                                                                                                                                            expression in developing seeds and so may be used
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 uptake activity in developing seeds Disclosure; Page 34~36; 58pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Glycine max.
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                                                                                                                                                                                                                                   trånsgene in developing seeds.
Sequence 524 AA;
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22-MAY-1997; US-047568.
(UNIW ) UNIV WASHINGTON STATE RES FOUND
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                                                                                                                                                                                                                    129 CQKHCQRRETEPR-HMQTCQQRCERRYEKEKRKQQKRYEEQQREDEEKYEERMKEED
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                                                                                                                                                                                                                                                                                 Local Similarity 31.6%; nes 18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity
les 18; Conse
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                                                                                                                                                      standard;
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larity 31.6%;
Conservative
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                                                                                                                                                      Protein;
                                                                                                                                                                                                                                                                                              23.9%;
                                                                                                                                                        593
                                                                                                                                                                                                                                                                             Score 129; DB 1; Len
Pred. No. 1.90e-02;
Pred. "----+ches 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 129; DB 1; L
Pred. No. 1.90e-02;
16; Mismatches 20;
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regions confer specific or enhanced
and so may be used to express any
                                                                                                                                                                                                                                                                                                              Length 524
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Best Local Similarity
Thehes 21; Conse
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
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                                                                                                                                                                                                         21-MAY-1998; U10465.
22-MAY-1997; US-047568.
(UNIW ) UNIV WASHINGTON S
Chao WS, Grimes HD;
WPI; 99-070155/06.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Novel anti-microbial protein from e.g. Macadamia integrifol useful for controlling microbial infestations of plants or Claim 1; Page 58-60; 96pp; English.

The sequence is that of an antimicrobial protein which can be used to control microbial infestations in plants and man
from Glycine max. This protein is used in a method resulting in the production of a modified plant sucrose binding protein (SBP) which has modified amino acid sequence compared to a corresponding wild-type SBP.
                                                   Claim 7; Page 39-40; 58pp; English.
This sequence represents a novel sucrose binding from Glycine max. This protein is used in a metho
                                                                                                                              New modified plant sucrose binding proteins - used to develop transgenic plants which can have enhanced or decreased sucrosuptake activity in developing seeds
                                                                                                                                                                                                                                                                                                                                                                                                   Glycine max.
                                                                                                                                                                                                                                                                                                                                                                                                                   SBP1; sucrose bindi
seed; carbohydrate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               animals.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       useful for controlling microbial infestations of plants or Disclosure; Page 60; 96pp; English.
The sequence is that of an antimicrobial protein which can
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Novel anti-microbial protein from e.g. Macadamia integrifolia useful for controlling microbial infestations of plants or mar
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (RETR-) COOP RES CENT TROPICAL PLANT PATHOLOGY Bower NI, Goulter KC, Green JL, Manners JM, MawPI; 98-377279/32.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      02-JUL-1998.
22-DEC-1997; AU0874
20-DEC-1996; AU-004
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                                                                                                                                                                                                                                                                                                                                                                     WO9853086-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    W90342 standard; protein; 409
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    be used to control microbial infestations in plants and mammalian
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO9827805-A1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Bower NI,
WPI; 98-3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             27-OCT-1998
                                                                                                                                                                                                                                                                                                                                             26-NOV-1998.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          128 QCQKHCQRR-ETEPRHMQTCQQRCERR 153
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              132 HCQRR-ETEPRHMQT-CQQRCERRYEKEKRKQQKRYEEQQREDEEKYE-ERMKEEDN 185
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local
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                                                                                                                                                                                                                                                                                                                                                                                                              max truncated SBP2 protein.
P1; sucrose binding protein; SBP2; sucrose uptake;
ed; carbohydrate content; soybean.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        6 ECRRQCLRRHEGQPWETQECMRRCRRR 32 : | : : : | | | | : | | | | | | |
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  mays antimicrobial
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
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Pred. No. 6.38e-02
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1. No. 2.98e-01;
1. No. ---hes 7;
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R26941 standard; Protein; R26941; 08-FEB-1993 (first entry)

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Best Local S
Matches 2
                                                                                                                                                                                                                                                                                                                   The products of the invention can be used for producing transgenic plants which have modified sucrose uptake activity, particularly in developing seeds. Enhanced sucrose uptake activity in developing seeds may be desirable where it is an advantage to increase the carbohydrate content of the seed (e.g. where the seed is the primary plant material harvested, such as soybean). In contrast, decreased sucrose uptake activity in seeds might be desirable where the vegetative material of the plant is harvested. The SBP regulatory regions confer specific or enhanced expression in developing seeds and so may be used to express any
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              desirable where it is an advantage to increase the carbohydrate content of the seed (e.g. where the seed is the primary plant material harvested, such as soybean). In contrast, decreased sucrose uptake activity in seeds might be desirable where the vegetative material of the plant is harvested. The SBP regulatory regions confer specific or enhanced
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         and where expression of the modified SBP in a yeast assay system confers enhanced sucrose uptake compared to the corresponding wild-type SBP. The products of the invention can be used for producing transgenic plants which have modified sucrose uptake activity, particularly in developing seeds. Enhanced sucrose uptake activity in developing seeds may be
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   This sequence represents a novel sucrose binding protein, SBP2 isolated from Glycine max. This protein is used in a method resulting in the production of a modified plant sucrose binding protein (SBP) which has a modified amino acid sequence compared to a corresponding wild-type SBP, and where expression of the modified SBP in a yeast assay system confers enhanced sucrose uptake compared to the corresponding wild-type SBP.
                                                                                                                                                                                                                                                                 Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 uptake activity in developing seeds Claim 13b; Page 37-38; 58pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New modified plant sucrose binding proteins - used to develop transgenic plants which can have enhanced or decreased sucrose
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Glycine max
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21-MAY-1998;
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CQKHCQR-RETEPRHMQTCQQRCERRYEKEKRKQQKRYEEQQREDEEKYEERMKEEDN
                                                                                                                                                               Similarity
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Pred. No. 2.98e-01;
15; Mismatches 18;
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                                                                                                                                                               113; DB 1;
No. 2.98e-01;
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05-FEB-1992; F00104.
05-FEB-1991; FR-001286.
05-FEB 1991; FR-001286.
(INSP) INST PASTEUR.
Druilhe P, Guerin-Marcha
Polypeptide(s) derived from liver stage of plasmodium falcipa-
for vaccination against, treatment of and diagnosis of mala
Claim 2; Fig 8-10; 81pp; French.
The 3' part of the P. falciparum liver-stage specific antigen
gene codes for a polypeptide sequence which carries a T cell
characteristic of a protein produced in hepatocytes infected
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          lambda gtl1 was used to transform E.coli. The expression library was screened with human antisera against antigens of all stages of P. falciparum. The library was rescreened with antibodies affinity-purified on a clone which was able to recognise antibodies specific to the hepatic phase. About 40 clones were detected which produced a characteristic LSA epitope. The clone with the largest insert (950 bases) encoded LSA-R-NR containing a 12-repeat region followed by a non-repeat region. Preferred antigenic polypeptides of the invention are derived from the amino acid sequence of LSA-R-NR.
                                                                                                                                         WPI; 92-299985/36.
N-PSDB; Q28119.
Polypeptide(s) derived from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             P.falciparum LSA gene C-terminal region.
Malaria; hepatocyte; sporozoite; T-cell epitope; B-cell epitope;
paludism; liver stage-specific antigen.
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Disclosure; Fig 1; 8lpp; French.
A genomic DNA bank of Pifalciparum EcoRI fragments prepared in
lambda gtll was used to transform E.coli. The expression librar
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (INSP ) INST PASTEUR. Druilhe P, Guerin-Mar WPI; 92-299985/36.
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20-AUG-1992.
05-FEB-1992; F00104.
05-FEB-1991; FR-001286.
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214. .49
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/note= "contains 12 x 17mer repeats"
210. .316
/label= non-repeat_region
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treatment of and diagnosis of malaria
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Pred. No. 9.71e-01;
13; Mismatches 30;
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Bower NI, Goulter KC, Green JL, Manners JM, Marcus JP;
WPI; 98-377279/32.
Novel anti-microbial protein from e.g. Macadamia integri
useful for controlling microbial infestations of plants
Claim 1; Page 66; 96pp; English.
The sequence is that of an antimicrobial protein which controlling microbial infestations in plants and
be used to control microbial infestations in plants and
                                                                                                            Sequence
                                                                                                                                                                                                                                                                                                      Stenocarpus sinuatus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                vaccines against malaria.
Sequence 493 AA;
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20-DEC-1996; AU-004275.
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                                                                                                                                                                                                                                                                                                                    antimicrobial protein; infestation;
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                                                                                                                            animals
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